

L10 ANSWER 8 OF 24 MEDLINE  
 AN 95157182 MEDLINE  
 DN 95157182 PubMed ID: 7854056  
 TI Expression of the growth-associated protein B-50/GAP43 via a defective herpes-simplex virus vector results in profound morphological changes in **non-neuronal** cells.  
 AU Verhaagen J; Hermens W T; Oestreicher A B; Gispen W H; Rabkin S D; Pfaff D  
 W; Kaplitt M G  
 CS Rudolf Magnus Institute, Department of Medical Pharmacology, Utrecht, The Netherlands.  
 SO BRAIN RESEARCH. MOLECULAR BRAIN RESEARCH, (1994 Oct) 26 (1-2) 26-36.  
 Journal code: MBR; 8908640. ISSN: 0169-328X.  
 CY Netherlands  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199503  
 ED Entered STN: 19950322  
 Last Updated on STN: 19980206  
 Entered Medline: 19950316  
 AB This study describes the creation and application of a defective herpes simplex viral (HSV) vector for B-50/GAP-43, a neural growth-associated phosphoprotein. We demonstrate abundant expression of B-50/GAP-43 in cultured **non-neuronal** cells (African green monkey kidney cells [vero cells] and Rabbit skin cells) via this HSV vector.  
 When  
 B-50/GAP-43 was expressed in **non-neuronal** cells major morphological changes occurred that included **extensive** membrane ruffling, the formation of filopodia and long thin **extensions** reminiscent of neurites. These **extensions** often terminated in **growth cone**-like structures. Quantitation of these morphological changes at different times following infection demonstrates that the surface area of the B-50/GAP-43-expressing cells started to increase between 6 and 10 h post-infection. At 72 h, B-50/GAP-43-positive cells were 3.0 times larger in size and one third of the cells expressed long processes with a mean length of 165 +/- 14.5 microns.  
 Ultrastructural  
 studies of cells 48 h after infection revealed that B-50/GAP-43 is predominantly localized at the plasma membrane of the elaborated processes. Some immunoreactivity was associated with vesicular structures that appear to be in-transit in the processes. These observations suggest that B-50/GAP-43 acts at the plasmamembrane to induce a neuron-like morphology in **non-neuronal** cells persisting for several days in culture. In the future the defective viral vector will enable gene transfer to express B-50/GAP-43 in neurons in vivo in order  
 to  
 study its involvement in regenerative sprouting and neuroplasticity.

L10 ANSWER 10 OF 24 MEDLINE  
 AN 94230600 MEDLINE  
 DN 94230600 PubMed ID: 8175908  
 TI An amino-terminal domain of the growth-associated protein GAP-43 mediates its effects on filopodial formation and cell spreading.  
 AU Strittmatter S M; Valenzuela D; Fishman M C  
 CS Developmental Biology Laboratory, Massachusetts General Hospital, Charlestown.  
 SO JOURNAL OF CELL SCIENCE, (1994 Jan) 107 ( Pt 1) 195-204.  
 Journal code: HNK; 0052457. ISSN: 0021-9533.  
 CY ENGLAND: United Kingdom  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199406  
 ED Entered STN: 19940620  
 Last Updated on STN: 19980206  
 Entered Medline: 19940606  
 AB GAP-43 is a neuronal protein that is believed to be important to neuronal growth and nerve terminal plasticity. It is enriched on the inner surface of **growth cone** membranes, a localization that may depend upon palmitoylation of Cys3 and Cys4. It is a major substrate for protein kinase C, which phosphorylates Ser41. Isolated GAP-43 can bind to actin and to calmodulin, and can activate the heterotrimeric GTP-binding proteins, G(o) and Gi. A peptide consisting of the GAP-43 sequence 39-55 binds calmodulin, and an amino-terminal GAP-43 (1-10) peptide activates G(o), suggesting that these stretches may be functional domains of the intact protein. When expressed in **non-neuronal** cells, GAP-43 enhances filopodial **extension** and has effects upon cell spreading. We have examined the effects of various GAP-43 domains upon this assay, by expression of GAP-43, GAP-43 mutant proteins, and GAP-43-CAT fusion proteins in COS-7 cells. We find that the amino terminus (Met-Leu-Cys-Cys-Met-Arg-Arg-Thr-Lys-Gln) is an important contributor to these effects on cell shape. A GAP-43 protein mutant in Cys3 and Cys4 does not bind to the membrane, and is inactive. Mutants in Arg6 or Lys9 also are inactive, although they remain localized to particulate fractions; Arg7 mutants are active. A chimeric gene consisting of GAP-43 (1-10) fused to chloramphenicol acetyl transferase (CAT) also causes cell shape changes. As for GAP-43, the effects of this fusion protein are abolished by mutations of Cys3, Cys4, Arg6 or Lys9, but not by mutation of Arg7. Therefore, the cell surface activity of transfected GAP-43 depends upon its amino terminus, although other domains may regulate it in this regard. Since the amino-terminal domain includes the peptide stretch known to be capable of activating G(o) and Gi, we examined the effect of GAP-43 on a Gi-regulated second messenger system, the inhibition of cAMP production in A431 cells. A431 cells stably transfected with GAP-43 spread less well than do controls. In addition, they evidence decreased levels of forskolin-stimulated cAMP, consistent with chronic stimulation of Gi. Stimulation of adenylate cyclase by isoproterenol reverses the GAP-43-induced changes in cell shape. This suggests that G protein stimulation is involved in GAP-43 effects upon cell shape.

L14 ANSWER 1 OF 174 MEDLINE  
 AN 1999028889 MEDLINE  
 DN 99028889 PubMed ID: 9812346  
 TI Effect of sublethal doses of cadmium, inorganic mercury and methylmercury on the cell morphology of an insect cell line (Aedes albopictus, C6/36).  
 AU Braeckman B; Simoens C; Rzeznik U; Raes H  
 CS Department of Biochemistry, Physiology and Microbiology, University of Gent, Belgium.  
 SO CELL BIOLOGY INTERNATIONAL, (1997 Dec) 21 (12) 823-32.  
 Journal code: BPN; 9307129. ISSN: 1065-6995.  
 CY ENGLAND: United Kingdom  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199812  
 ED Entered STN: 19990115  
 Last Updated on STN: 19990115  
 Entered Medline: 19981201  
 AB The effect of CdCl<sub>2</sub> (44 microM), HgCl<sub>2</sub> (3.7 microM), and MeHgCl (2 microM) on the morphology of Aedes albopictus C6/36 cells was studied at the light microscopical level. Treatment times and metal concentrations were in the sublethal range as determined by a fluorometric dye exclusion test. The three metal species had profound effects on the cell morphology. MeHgCl treatment induced the development of a large number of short, actin-supported, tangled **filopodia**. Both CdCl<sub>2</sub> and HgCl<sub>2</sub> induced long **extensions**. Pretreatment with colchicine but not with cytochalasin B prevented formation of these **extensions** which suggests that they were supported by microtubules. This was confirmed by immunostaining for microtubules. The extensions were relatively stable towards colchicine post-treatment. To authors' knowledge, this effect has not yet been described for heavy metals. The similarity with 20-hydroxyecdysone-treated cells and the occurrence of cytoplasmic feet in insect cells is discussed.

L14 ANSWER 4 OF 174 MEDLINE  
 AN 1998028160 MEDLINE  
 DN 98028160 PubMed ID: 9362067  
 TI Integrin alpha 6A beta 1 induces CD81-dependent cell motility without engaging the extracellular matrix migration substrate.  
 AU Domanico S Z; Pelletier A J; Havran W L; Quaranta V  
 CS Department of Cell Biology, Scripps Research Institute, La Jolla, California 92037, USA.  
 NC CA-47858 (NCI)  
 DE-10063 (NIDCR)  
 SO MOLECULAR BIOLOGY OF THE CELL, (1997 Nov) 8 (11) 2253-65.  
 Journal code: BAU; 9201390. ISSN: 1059-1524.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199712  
 ED Entered STN: 19980109  
 Last Updated on STN: 20000303  
 Entered Medline: 19971211  
 AB It is well established that integrins and extracellular matrix (ECM) play key roles in cell migration, but the underlying mechanisms are poorly defined. We describe a novel mechanism whereby the integrin alpha 6 beta 1, a laminin receptor, can affect cell motility and induce migration onto ECM substrates with which it is not engaged. By using DNA-mediated gene transfer, we expressed the human integrin subunit alpha 6A in murine embryonic stem (ES) cells. ES cells expressing alpha 6A (ES6A) at the surface dimerized with endogenous beta 1, **extended** numerous **filopodia** and lamellipodia, and were intensely migratory in haptotactic assays on laminin (LN)-1. Transfected alpha 6A was responsible for these effects, because cells transfected with control vector or alpha 6B, a cytoplasmic domain alpha 6 isoform, displayed compact morphology and no migration, like wild-type ES cells. The ES6A migratory phenotype persisted on fibronectin (Fn) and Ln-5. Adhesion inhibition assays indicated that alpha 6 beta 1 did not contribute detectably to adhesion to these substrates in ES cells. However, anti-alpha 6 antibodies completely blocked migration of ES6A cells on Fn or Ln-5. Control experiments with monensin and anti-ECM antibodies indicated that this inhibition could not be explained by deposition of an alpha 6 beta 1 ligand (e.g., Ln-1) by ES cells. Cross-linking with secondary antibody overcame the inhibitory effect of anti-alpha 6 antibodies, restoring migration or **filopodia extension** on Fn and Ln-5. Thus, to induce migration in ES cells, alpha 6A beta 1 did not have to engage with an ECM ligand but likely participated in molecular interactions sensitive to anti-alpha 6 beta 1 antibody and mimicked by cross-linking. Antibodies to the tetraspanin CD81 inhibited alpha 6A beta 1-induced migration but had no effect on ES cell adhesion. It is known that CD81 is physically associated with alpha 6 beta 1, therefore our results suggest a mechanism by which interactions between alpha 6A beta 1 and CD81 may up-regulate cell motility, affecting migration mediated by other integrins.

L14 ANSWER 5 OF 174 MEDLINE  
 AN 97412181 MEDLINE  
 DN 97412181 PubMed ID: 9265651  
 TI PSTPIP: a tyrosine phosphorylated cleavage furrow-associated protein that is a substrate for a PEST tyrosine phosphatase.

AU Spencer S; Dowbenko D; Cheng J; Li W; Brush J; Utzig S; Simanis V; Lasky  
L  
A  
CS Department of Molecular Oncology, Genentech, Inc., South San Francisco,  
California 94080, USA.  
SO JOURNAL OF CELL BIOLOGY, (1997 Aug 25) 138 (4) 845-60.  
Journal code: HMV; 0375356. ISSN: 0021-9525.  
CY United States  
DT Journal; Article; (JOURNAL ARTICLE)  
LA English  
FS Priority Journals  
OS GENBANK-U87814  
EM 199709  
ED Entered STN: 19970926  
Last Updated on STN: 19990129  
Entered Medline: 19970915  
AB We have investigated proteins which interact with the PEST-type protein  
tyrosine phosphatase, PTP hematopoietic stem cell fraction (HSCF), using  
the yeast two-hybrid system. This resulted in the identification of  
proline, serine, threonine phosphatase interacting protein (PSTPIP), a  
novel member of the actin- associated protein family that is homologous  
to  
Schizosaccharomyces pombe CDC15p, a phosphorylated protein involved with  
the assembly of the actin ring in the cytokinetic cleavage furrow. The  
binding of PTP HSCF to PSTPIP was induced by a novel interaction between  
the putative coiled-coil region of PSTPIP and the COOH-terminal,  
proline-rich region of the phosphatase. PSTPIP is tyrosine phosphorylated  
both endogenously and in v-Src transfected COS cells, and cotransfection  
of dominant-negative PTP HSCF results in hyperphosphorylation of PSTPIP.  
This dominant-negative effect is dependent upon the inclusion of the  
COOH-terminal, proline-rich PSTPIP-binding region of the phosphatase.  
Confocal microscopy analysis of endogenous PSTPIP revealed colocalization  
with the cortical actin cytoskeleton, lamellipodia, and actin-rich  
cytokinetic cleavage furrow. Overexpression of PSTPIP in 3T3 cells  
resulted in the formation of **extended filopodia**,  
consistent with a role for this protein in actin reorganization. Finally,  
overexpression of mammalian PSTPIP in exponentially growing S. pombe  
results in a dominant-negative inhibition of cytokinesis. PSTPIP is  
therefore a novel actin-associated protein, potentially involved with  
cytokinesis, whose tyrosine phosphorylation is regulated by PTP HSCF.

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L14 ANSWER 2 OF 174 MEDLINE  
 AN 1998075097 MEDLINE  
 DN 98075097 PubMed ID: 9412479  
 TI The integrin alpha6beta4 functions in carcinoma cell migration on laminin-1 by mediating the formation and stabilization of actin-containing motility structures.  
 AU Rabinovitz I; Mercurio A M  
 CS Department of Medicine (GI Division), Beth Israel Deaconess Medical Center and Harvard Medical School, Boston, Massachusetts 02215, USA.  
 NC AI39264 (NIAID)  
 CA44704 (NCI)  
 SO JOURNAL OF CELL BIOLOGY, (1997 Dec 29) 139 (7) 1873-84.  
 Journal code: HMV; 0375356. ISSN: 0021-9525.  
 CY United States  
 DT Journal; Article; (JOURNAL ARTICLE)  
 LA English  
 FS Priority Journals  
 EM 199802  
 ED Entered STN: 19980217  
 Last Updated on STN: 20000303  
 Entered Medline: 19980205  
 AB Functional studies on the alpha6beta4 integrin have focused primarily on its role in the organization of hemidesmosomes, stable adhesive structures that associate with the intermediate filament cytoskeleton. In this study, we examined the function of the alpha6beta4 integrin in clone A cells, a colon carcinoma cell line that expresses alpha6beta4 but no alpha6beta1 integrin and exhibits dynamic adhesion and motility on laminin-1. Time-lapse videomicroscopy of clone A cells on laminin-1 revealed that their migration is characterized by **filopodial extension** and stabilization followed by lamellae that **extend** in the direction of stabilized **filopodia**. A function-blocking mAb specific for the alpha6beta4 integrin inhibited clone A migration on laminin-1. This mAb also inhibited filopodial formation and stabilization and lamella formation. Indirect immunofluorescence microscopy revealed that the alpha6beta4 integrin is localized as discrete clusters in filopodia, lamellae, and retraction fibers. Although beta1 integrins were also localized in the same structures, a spatial separation of these two integrin populations was evident. In filopodia and lamellae, a striking colocalization of the alpha6beta4 integrin and F-actin was seen. An association between alpha6beta4 and F-actin is supported by the fact that alpha6beta4 integrin and actin were released from clone A cells by treatment with the F-actin-severing protein gelsolin and that alpha6beta4 immunostaining at the marginal edges of clone A cells on laminin-1 was resistant to solubilization with Triton X-100. Cytokeratins were not observed in filopodia and lamellipodia. Moreover, alpha6beta4 was extracted from these marginal edges with a Tween-40/deoxycholate buffer that solubilizes the actin cytoskeleton but not cytokeratins. Three other carcinoma cell lines (MIP-101, CCL-228, and MDA-MB-231) exhibited alpha6beta4 colocalized with actin in filopodia and lamellae. Formation of lamellae in these cells was inhibited with an alpha6-specific antibody. Together, these results indicate that the alpha6beta4 integrin functions in carcinoma migration on laminin-1 through its ability to promote the formation and stabilization of actin-containing motility structures.

L2 ANSWER 1 OF 1 MEDLINE  
AN 1999223289 MEDLINE  
DN 99223289 PubMed ID: 10208590  
TI **Isolation of human delta-catenin**  
and its binding specificity with presenilin 1.  
AU Tanahashi H; Tabira T  
CS Division of Demyelinating Disease and Aging, National Institute of  
Neuroscience, Kodaira, Tokyo, Japan.  
SO NEUROREPORT, (1999 Feb 25) 10 (3) 563-8.  
Journal code: 9100935. ISSN: 0959-4965.  
CY ENGLAND: United Kingdom  
DT Journal; Article; (JOURNAL ARTICLE)  
LA English  
FS Priority Journals  
OS GENBANK-AB013805  
EM 199906  
ED Entered STN: 19990618  
Last Updated on STN: 19990618  
Entered Medline: 19990610  
AB We screened proteins for interaction with presenilin (PS) 1, and cloned  
the full-length cDNA of human delta-catenin, which encoded 1225 amino  
acids. Yeast two-hybrid assay, GST binding assay and immunoprecipitation  
demonstrated that delta-catenin interacted with a hydrophilic loop region  
in the endoproteolytic C-terminal fragment of PS1, but not with that of  
PS-2. These results suggest that PS1 and PS2 partly differ in function.  
PS1 loop fragment containing the pathogenic mutation retained the binding  
ability. We also found another armadillo-protein, p0071, interacted with  
PS1.